

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/053,498c
Source: IFW16
Date Processed by STIC: 01/05/2006

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 01/05/2006

PATENT APPLICATION: US/10/053,498C

TIME: 14:32:32

Input Set : E:\8449-429-999 (substitute seqlisting).txt

Output Set: N:\CRF4\01052006\J053498C.raw

3 <110> APPLICANT: Sloan-Kettering Institute for Cancer Research
 4 Rothman, James E.
 5 Mayhew, Mark
 6 Hoe, Mee H.
 7 Houghton, Alan
 8 Hartl, Ulrich
 9 Ouerfelli, Ouathak
 10 Moroi, Yoichi
 12 <120> TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
 14 <130> FILE REFERENCE: 8449-429-999
 16 <140> CURRENT APPLICATION NUMBER: 10/053,498C
 17 <141> CURRENT FILING DATE: 2002-01-17
 19 <150> PRIOR APPLICATION NUMBER: 08/961,707
 20 <151> PRIOR FILING DATE: 1997-10-31
 22 <160> NUMBER OF SEQ ID NOS: 326
 24 <170> SOFTWARE: PatentIn 3.0
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 8
 28 <212> TYPE: PRT
 29 <213> ORGANISM: Artificial Sequence
 31 <220> FEATURE:
 32 <223> OTHER INFORMATION: peptide in m13 coliphage
 34 <400> SEQUENCE: 1
 36 His Thr Thr Val Tyr Gly Ala Gly
 37 1 5
 39 <210> SEQ ID NO: 2
 40 <211> LENGTH: 8
 41 <212> TYPE: PRT
 42 <213> ORGANISM: Artificial Sequence
 44 <220> FEATURE:
 45 <223> OTHER INFORMATION: peptide in m13 coliphage
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 49 Thr Glu Thr Pro Tyr Pro Thr Gly
 50 1 5
 52 <210> SEQ ID NO: 3
 53 <211> LENGTH: 8
 54 <212> TYPE: PRT
 55 <213> ORGANISM: Artificial Sequence
 57 <220> FEATURE:
 58 <223> OTHER INFORMATION: peptide in m13 coliphage
 60 <400> SEQUENCE: 3
 62 Leu Thr Thr Pro Phe Ser Ser Gly
 63 1 5

Cp9-67

RAW SEQUENCE LISTING

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Input Set : E:\8449-429-999 (substitute seqlisting).txt

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67 <212> TYPE: PRT
68 <213> ORGANISM: Artificial Sequence
70 <220> FEATURE:
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73 <400> SEQUENCE: 4
75 Gly Val Pro Leu Thr Met Asp Gly
76 1 5
78 <210> SEQ ID NO: 5
79 <211> LENGTH: 8
80 <212> TYPE: PRT
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: peptide in m13 coliphage
86 <400> SEQUENCE: 5
88 Lys Leu Pro Thr Val Leu Arg Gly
89 1 5
91 <210> SEQ ID NO: 6
92 <211> LENGTH: 8
93 <212> TYPE: PRT
94 <213> ORGANISM: Artificial Sequence
96 <220> FEATURE:
97 <223> OTHER INFORMATION: peptide in m13 coliphage
99 <400> SEQUENCE: 6
101 Cys Arg Phe His Gly Asn Arg Gly
102 1 5
104 <210> SEQ ID NO: 7
105 <211> LENGTH: 8
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: peptide in m13 coliphage
112 <400> SEQUENCE: 7
114 Tyr Thr Arg Asp Phe Glu Ala Gly
115 1 5
117 <210> SEQ ID NO: 8
118 <211> LENGTH: 8
119 <212> TYPE: PRT
120 <213> ORGANISM: Artificial Sequence
122 <220> FEATURE:
123 <223> OTHER INFORMATION: peptide in m13 coliphage
125 <400> SEQUENCE: 8
127 Ser Ser Ala Ala Gly Pro Arg Gly
128 1 5
130 <210> SEQ ID NO: 9
131 <211> LENGTH: 8
132 <212> TYPE: PRT
133 <213> ORGANISM: Artificial Sequence

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136 <223> OTHER INFORMATION: peptide in m13 coliphage
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141     1               5
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146 <213> ORGANISM: Artificial Sequence
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149 <223> OTHER INFORMATION: peptide in m13 coliphage
151 <220> FEATURE:
152 <221> NAME/KEY: MISC_FEATURE
153 <222> LOCATION: (7)..(7)
154 <223> OTHER INFORMATION: unknown
156 <400> SEQUENCE: 10
W--> 158   Asp Ala Leu Met Trp Pro Xaa Gly
159     1               5
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163 <212> TYPE: PRT
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: peptide in m13 coliphage
169 <220> FEATURE:
170 <221> NAME/KEY: MISC_FEATURE
171 <222> LOCATION: (3)..(3)
172 <223> OTHER INFORMATION: unknown
174 <400> SEQUENCE: 11
W--> 176   Ser Ser Xaa Ser Leu Tyr Ile Gly
177     1               5
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184 <220> FEATURE:
185 <223> OTHER INFORMATION: peptide in m13 coliphage
187 <400> SEQUENCE: 12
189   Phe Asn Thr Ser Thr Arg Thr Gly
190     1               5
192 <210> SEQ ID NO: 13
193 <211> LENGTH: 8
194 <212> TYPE: PRT
195 <213> ORGANISM: Artificial Sequence
197 <220> FEATURE:
198 <223> OTHER INFORMATION: peptide in m13 coliphage
200 <400> SEQUENCE: 13
202   Thr Val Gln His Val Ala Phe Gly
203     1               5

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216       1           5
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219 <211> LENGTH: 8
220 <212> TYPE: PRT
221 <213> ORGANISM: Artificial Sequence
223 <220> FEATURE:
224 <223> OTHER INFORMATION: peptide in m13 coliphage
226 <400> SEQUENCE: 15
228   Val Gly Ser Met Glu Ser Leu Gly
229       1           5
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233 <212> TYPE: PRT
234 <213> ORGANISM: Artificial Sequence
236 <220> FEATURE:
237 <223> OTHER INFORMATION: peptide in m13 coliphage
239 <220> FEATURE:
240 <221> NAME/KEY: MISC_FEATURE
241 <222> LOCATION: (2)..(2)
242 <223> OTHER INFORMATION: unknown
244 <220> FEATURE:
245 <221> NAME/KEY: MISC_FEATURE
246 <222> LOCATION: (6)..(6)
247 <223> OTHER INFORMATION: unknown
249 <400> SEQUENCE: 16
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252       1           5
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255 <211> LENGTH: 8
256 <212> TYPE: PRT
257 <213> ORGANISM: Artificial Sequence
259 <220> FEATURE:
260 <223> OTHER INFORMATION: peptide in m13 coliphage
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264   Ala Pro Pro Arg Val Thr Met Gly
265       1           5
267 <210> SEQ ID NO: 18
268 <211> LENGTH: 8
269 <212> TYPE: PRT
270 <213> ORGANISM: Artificial Sequence
272 <220> FEATURE:

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273 <223> OTHER INFORMATION: peptide in m13 coliphage
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277 Ile Ala Thr Lys Thr Pro Lys Gly
278 1 5
280 <210> SEQ ID NO: 19
281 <211> LENGTH: 8
282 <212> TYPE: PRT
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
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288 <400> SEQUENCE: 19
290 Lys Pro Pro Leu Phe Gln Ile Gly
291 1 5
293 <210> SEQ ID NO: 20
294 <211> LENGTH: 8
295 <212> TYPE: PRT
296 <213> ORGANISM: Artificial Sequence
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301 <400> SEQUENCE: 20
303 Tyr His Thr Ala His Asn Met Gly
304 1 5
306 <210> SEQ ID NO: 21
307 <211> LENGTH: 8
308 <212> TYPE: PRT
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
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314 <400> SEQUENCE: 21
316 Ser Tyr Ile Gln Ala Thr His Gly
317 1 5
319 <210> SEQ ID NO: 22
320 <211> LENGTH: 8
321 <212> TYPE: PRT
322 <213> ORGANISM: Artificial Sequence
324 <220> FEATURE:
325 <223> OTHER INFORMATION: peptide in m13 coliphage
327 <400> SEQUENCE: 22
329 Ser Ser Phe Ala Thr Phe Leu Gly
330 1 5
332 <210> SEQ ID NO: 23
333 <211> LENGTH: 8
334 <212> TYPE: PRT
335 <213> ORGANISM: Artificial Sequence
337 <220> FEATURE:
338 <223> OTHER INFORMATION: peptide in m13 coliphage
340 <400> SEQUENCE: 23
342 Thr Thr Pro Pro Asn Phe Ala Gly
343 1 5

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/053,498C

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Input Set : E:\8449-429-999 (substitute seqlisting).txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:10; Xaa Pos. 7
Seq#:11; Xaa Pos. 3
Seq#:16; Xaa Pos. 2,6
Seq#:47; N Pos. 19
Seq#:48; N Pos. 8
Seq#:53; N Pos. 6,16,18
Seq#:100; Xaa Pos. 5
Seq#:133; N Pos. 14
Seq#:251; Xaa Pos. 7
Seq#:252; Xaa Pos. 3
Seq#:257; Xaa Pos. 2,6
Seq#:304; Xaa Pos. 5
Seq#:322; Xaa Pos. 1,2,3,4,5
Seq#:323; Xaa Pos. 1,2,3,4
Seq#:324; Xaa Pos. 1,2,3,4,5,6
Seq#:325; Xaa Pos. 1,2,3,4
Seq#:326; Xaa Pos. 1,2,3,4,5,6,7

VERIFICATION SUMMARY

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Input Set : E:\8449-429-999 (substitute seqlisting).txt

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L:158 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:650 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:0
L:667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:0
L:742 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:0
L:1336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100 after pos.:0
L:1745 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:133 after pos.:0
L:3276 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:251 after pos.:0
L:3294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:252 after pos.:0
L:3369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:257 after pos.:0
L:3985 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:304 after pos.:0
L:4231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:322 after pos.:0
L:4255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:323 after pos.:0
L:4279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:324 after pos.:0
L:4303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:325 after pos.:0
L:4332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:326 after pos.:0